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ID
     Q41238 SOLTU PRELIMINARY;
                                     PRT:
                                            857 AA.
AC
DT
     01-NOV-1996 (TrEMBLrel. 01, Created)
DT
     01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
     01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DT
     Linoleate:oxygen oxidoreductase (Fragment).
DE
OS
     Solanum tuberosum (Potato).
OC
     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
OC
     Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
OC
     lamiids; Solanales; Solanaceae; Solanum.
OX
     NCBI TaxID=4113;
RN
RP
     NUCLEOTIDE SEQUENCE.
RX
     MEDLINE=94302170; PubMed=8029354; DOI=10.1104/pp.105.1.269;
RA
     Geerts A., Feltkamp D., Rosahl S.;
RT
     "Expression of lipoxygenase in wounded tubers of Solanum tuberosum
RT
RL
     Plant Physiol. 105:269-277(1994).
CC
     -!- FUNCTION: Plant lipoxygenase may be involved in a number of
CC
         diverse aspects of plant physiology including growth and
CC
         development, pest resistance, and senescence or responses to
CC
         wounding (By similarity).
CC
     -!- CATALYTIC ACTIVITY: Linoleate + O(2) = (92,11E)-(13S)-13-
CC
         hydroperoxyoctadeca-9,11-dienoate.
CC
     -!- COFACTOR: Iron (By similarity).
CC
     -!- SIMILARITY: Belongs to the lipoxygenase family.
CC
     -!- SIMILARITY: Contains 1 PLAT domain.
DR
     EMBL; <u>$73865</u>; <u>AAB31252</u>.1; -.
DR
     HSSP; P08170; 1FGT.
DR
     GO; GO: 0005506; F:iron ion binding; IEA.
DR
     GO; GO:0016165; F:lipoxygenase activity; IEA.
DR
     GO; GO:0016491; F:oxidoreductase activity; IEA.
     GO; GO:0006118; P:electron transport; IEA.
DR
DR
     InterPro; <u>IPR000907</u>; Lipoxygenase.
     InterPro; IPR001024; Lipoxygenase_LH2.
DR
     InterPro; <u>IPR001246</u>; Plant_lipoxygnse.
DR
     InterPro; IPR008976; PLAT LH2.
DR
     Pfam; PF00305; Lipoxygenase; 1.
DR
     Pfam; <u>PF01477</u>; <u>PLAT</u>; 1.
DR
     PRINTS; PR00087; LIPOXYGENASE.
DR
     PRINTS; PR00468; PLTLPOXGNASE.
DR
     SMART; SM00308; LH2; 1.
DR
     PROSITE; PS00711; LIPOXYGENASE 1; 1.
DR
     PROSITE; PS00081; LIPOXYGENASE 2; 1.
     PROSITE; PS50095; PLAT; 1.
KW
     Dioxygenase; Iron; Oxidoreductase.
     NON TER
FT
                 857
                        857
SQ
     SEQUENCE
                857 AA; 96585 MW; 3785A24E8DBA8DA7 CRC64;
    QIVGGLIGGH HDSKKVKGTV VMMKKNALDF TDLAGSLTDK IFEALGQKVS FQLISSVQSD
  PANGLQGKHS NPAYLENFLF TLTPLAAGET AFGVTFDWNE EFGVPGAFII KNTHINEFFL
  W KSLTLEDVPN HGKVHFVCNS WVYPSFRYKS DRIFFANQPY LPSETPELLR KYRENELLTL
  I RGDGTGKREA WDRIYDYDVY NDLGNPDQGE QNVRTTLGGS ADYPYPRRGR TGRPPTRTDP
  A KSESRIPLIL SLDIYVPRDE RFGHLKMSDF LTYALKSIVQ FILPELHALF DGTPNEFDSF
  EDVLRLYEGG IKLPQGPLFK ALTAAIPLEM MKELLRTDGE GILRFPTPLV IKDSKTAWRT
  → DEEFAREMLA GVNPIIISRL QEFPPKSKLD PEAYGNQNST ITAEHIEDKL DGLTVDEAMN
 NNKLFILNHH DVLIPYLRRI NTTTTKTYAS RTLLFLQDNG SLKPLAIELS LPHPDGDQFG
 VISKVYTPSD QGVESSIWQL AKAYVAVNDS GVHQLISHWL NTHAVIEPFV IATNRQLSVL
 WO PADLVKRGVA VEDSSSPHGV RLLIEDYPYA VDGLEIWSAI KSWVTDYCSF YYGSDEEILK
 DNELQAWWKE LREVGHGDKK NEPWWPEMET PQELIDSCTT IIWIASALHA AVNFGQYPYA
                                      171
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